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OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/553,094

DATE: 05/09/2000
TIME: 16:28:06

Input Set : D:\AprilCornReg.rpt
Output Set: N:\CRF3\05092000\I553094.raw

1 <110> APPLICANT: Andersen, Scott E.
 2 Fisher, Dane K.
 4 <120> TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
 5 Plants
 7 <130> FILE REFERENCE: 38-21(15502)B
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/553,094
 C--> 9 <141> CURRENT FILING DATE: 2004-04-18
 9 <160> NUMBER OF SEQ ID NOS: 18831
 11 <210> SEQ ID NO: 1
 12 <211> LENGTH: 341
 13 <212> TYPE: DNA
 14 <213> ORGANISM: Zea mays
 16 <223> OTHER INFORMATION: Clone ID: LIB3061-001-Q1-K2-H1
 18 <400> SEQUENCE: 1
 20 gtctctgcta ctcaccttagc cgccagtcat gggcatgatc aagatcgaa tcaacgggtt 60
 22 cggaggatc ggctggctcg tggcagggt cgtccctgcag agcgaggatg tcgactctgt 120
 24 ctgcgtcaac gaccgcgttca tcaccacggta ttacatgacc tacatgttca agtacgacac 180
 26 cgtgcacgggt caatggaaac acatgcgaca ttaccctcaa ggactccaag acgcttcttg 240
 28 ttggcggagaa gccggtcacc gcgtttggca tcaagaacccct ccggagatggccgtatgtg 300
 30 aggctgtgtc tgatatgtc gtgcgttca tcgggtgttcc 341
 33 <210> SEQ ID NO: 2
 34 <211> LENGTH: 371
 35 <212> TYPE: DNA
 36 <213> ORGANISM: Zea mays
 38 <223> OTHER INFORMATION: Clone ID: LIB3061-005-Q1-K1-B3
 40 <400> SEQUENCE: 2
 42 ctcccagttc gatccccatt taaaaaacac ccatccttcg ttccctccgtc tagcagccgc 60
 44 tcaccctcat ggccaaagatc aagatcgaaa tcaatgggtt tggcaggatc ggaaggctgg 120
 46 tggccagggt ggccctgcag agcgtacgacg tcgactctgt cggccgtaaac gacccttca 180
 48 tcagcacccga ttacatgacc tacatgttca agtacgacac tgcgtacggaa cagtggaaac 240
 50 accatggatgtt caagggtggaa gactccaaa cccttcttcc ttggtgagaaa gagggttgctg 300
 52 tggccagggtt ccgttccat gaggatgttcc catgggttcc ttggtgccgtt gaatatgtt 360
 54 ttgttgttac t 371
 57 <210> SEQ ID NO: 3
 58 <211> LENGTH: 328
 59 <212> TYPE: DNA
 60 <213> ORGANISM: Zea mays
 62 <223> OTHER INFORMATION: Clone ID: LIB3061-056-Q1-K1-F11
 64 <400> SEQUENCE: 3
 66 acggccgtcat cgatctcggt tcaaatccat ccatcctcggt ttccctccgtc ccgcagcatc 60
 68 tcaacccgtt gggaaagatc aagatcgaaa tcaatgggtt cggcaggatc ggaaggctgg 120
 70 tggccagggtt ggccctgcag agtacgatgt tcgactctgt cggccgtaaac gacccttca 180
 72 tcagcacccga ctacatgacc tacatgttca agtacgacac tgcgtacggaa cagtggaaac 240
 74 accatggatgtt caagggtggaa gactccaaa cccttcttcc ttggtgagaaa gagggttgctg 300
 76 tggccagggtt ccgttccat gaggatgttcc catgggttcc ttggtgccgtt gaatatgtt 328
 79 <210> SEQ ID NO: 4
 80 <211> LENGTH: 408

ENTERED

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81 <212> TYPE: DNA
82 <213> ORGANISM: Zea mays
84 <223> OTHER INFORMATION: Clone ID: LIB3061-013-Q1-K1-D10
86 <400> SEQUENCE: 4
88 cacagatcac acacactcgatctccatt tegatccccat ttgaaatcaa cccatcgctg 60
90 tgtccctcggt cttagcggct ctcaccctca tggccaagat caagatcggtt atcaatgggt 120
92 tcggcaggat cggaaaggctg gtggccagggt tgccctgcga gagegacgac gtcgagctcg 180
94 tcgcccgtgaa cgacccttc atcagcaccg attacatgac ctacatgttc aagtatgaca 240
96 ctgtgcacgg acatggaa caccatgggg tcaaggtaaa ggactccaag acccttctt 300
98 ttggtgagaa ggagggtcgct gtgtttggct gcaggaaccc tgaggagatc ccatggtct 360
100 cgggttgcgc tgaatatgtct gtcgagtcta ctgggtttt cactgacc 408
103 <210> SEQ ID NO: 5
104 <211> LENGTH: 417
105 <212> TYPE: DNA
106 <213> ORGANISM: Zea mays
108 <223> OTHER INFORMATION: Clone ID: LIB3061-011-Q1-K1-H1
110 <400> SEQUENCE: 5
112 cacactcgatccat cgtatccattt tgaaatcaac ccatcctcg ttcctccgtc 60
114 tagcagccctc tcaccctcat gggcaagatc aagatcggtt tggcaggatc 120
116 ggaaggctgg tggccagggt gggccctgcag aegcgaacgac tegagctcg tggccgtgaa 180
118 gacccttc tccatgcacc ttacatgacc tacatgttca agttagacatc tggcaggg 240
120 cagtggaaacccatcgggtt caatggtaag gactccaaga cccttctt tggtgagaa 300
122 gaagttgtcg tggccatggc cggggaccc tggaggatcc catgggggtc ggttggcgt 360
124 gaatatgttgc tggatctac tgggtttt actgaccagg agaaggctgc agtcac 417
127 <210> SEQ ID NO: 6
128 <211> LENGTH: 422
129 <212> TYPE: DNA
130 <213> ORGANISM: Zea mays
132 <223> OTHER INFORMATION: Clone ID: LIB3061-034-Q1-K1-H6
134 <400> SEQUENCE: 6
136 acacactcgatccat tccatctcat ttgaaatcaa cccatcctcg tggccctgggt 60
138 ctagcggct ctcaccctca tggccaagat caagatcggtt atcaatgggt tggcaggat 120
140 cggaaaggctg gtggccagggt tgccctgcga gagegacgac gtcgagctcg tggccgtgaa 180
142 cgacccttc atcagcaccg attacatgac ctacatgttc aagtatgaca ctgtgcacgg 240
144 acatggaaacccatcgggtt caatggtaaa ggactccaag acccttctt tggtgagaa 300
146 ggagggtcgct gtgtttggct gcaggaaccc tggaggatcc ccatgggggtc cgggtggcgc 360
148 tgaatatgttgc tggatctac tgggtttt actgaccagg gagaaggctg cagctcactt 420
150 ga 422
153 <210> SEQ ID NO: 7
154 <211> LENGTH: 420
155 <212> TYPE: DNA
156 <213> ORGANISM: Zea mays
158 <223> OTHER INFORMATION: Clone ID: LIB3061-011-Q1-K1-D5
160 <400> SEQUENCE: 7
162 tcggaaatcaa cggcttcggat aggatcggtt ggctcggtt cgggtcgcc ctccagacgc 60
164 aggtatcgat gtcgatcgcc gtcacacgacc ctttcatcac cacggactac atgacactaca 120
166 tggatcaatcgat cggacccgtt caccggactt ggaaggcacatc gacatcaca ctcaaggact 180
168 ccaagacccat tctttcggtt gacaaggccgg tccaccgtttt tggcatcagg aaccctgggg 240
170 aaatccctgtt ggggtggatc ggcgtcgat atgtcgatggat gtcacccggc gtcttcactg 300

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Input Set : D:\AprilCornReg.rpt
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 Output Set: N:\CRF3\05092000\I553094.raw

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260 tcgccccggg gccgcagggt accagccgcg ttggcttgc acacaagccc acaaacacag 120
262 ggcgcctccgc acggacaaaa tctccgaaac gcccattggct aaggaagaga gggttatcg 180
264 aagtgaata ttgacatggg attagcatcg acaagttttt attagactaa aagacacat 240
266 catatttggg cttttggaga gagcacagtt ttgttacaac gtcgtacat atgatagcaa 300
268 tgctttcac atggatggt ttggagggtc ttgggttggaa tatatgggtt gagaaactga 360
270 aaagctccat gcacagggtt ggaggtacaa gagcccagat gagcaccctt tctttccaa 420
272 ggatctgcct gagcc 435
275 <210> SEQ ID NO: 12
276 <211> LENGTH: 379
277 <212> TYPE: DNA
278 <213> ORGANISM: Zea mays
280 <223> OTHER INFORMATION: Clone ID: LIB3061-035-Q1-K1-B12
282 <400> SEQUENCE: 12
284 cctcccaactt cgtgcgagcg tccegaacta agttgtctgt ggtggagggtt gtttgtggcg 60
286 atggccctca agctggtcac caagccgcg gcccgtcgc ccgtctgc tcactgggg 120
288 gatctcgccc ggtggccca ggttacgacg cgcgttgccct tcggaccacg gcccaggAAC 180
290 aaggggctcc gcacggggca caactccgc acggccatgg ctaaggaga gagggttgat 240
292 cgaatgtaaa tattggacattt ggatagcata agacaatgtt tgatttagact agaagacacg 300
294 atcatatttgg gacttttggg gagagcacag tttgttaca acgctgacac atatgatagc 360
296 aatgctatcc acatggatg 379
299 <210> SEQ ID NO: 13
300 <211> LENGTH: 392
301 <212> TYPE: DNA
302 <213> ORGANISM: Zea mays
304 <223> OTHER INFORMATION: Clone ID: LIB3061-034-Q1-K1-D8
306 <400> SEQUENCE: 13
308 caaattccata ctttcaatca gagccagttt cctcgattt ttaccgggaa cttcttgc 60
310 ctteccagca aaggaaatc tggaaaggat ggctggatc ctcctggtca tggtgatgt 120
312 ttccctctt tgaataacag cgaaaactt gacatcttat tggctcaggaa caaaggat 180
314 gtctttgtt caaactcaga caacttgggtt gctatgtcg acatcaagat cctaaaccat 240
316 ctgatcataa accagaacga gtactgcattt gaagtttactt caaaagacgtt ggctgacgtt 300
318 aaaggtggca ctctccatctc ttacgaaagg aagatgttgc cttttggag attgccaag 360
320 tacccgatga gcatgtgaat gaatttaat ca 392
323 <210> SEQ ID NO: 14
324 <211> LENGTH: 369
325 <212> TYPE: DNA
326 <213> ORGANISM: Zea mays
328 <223> OTHER INFORMATION: unsure at all n locations
329 <223> OTHER INFORMATION: Clone ID: LIB3061-040-Q1-K1-B8
331 <400> SEQUENCE: 14
333 ctggctgaag tgcgtgcctt cttcgtatgtt catgaccaag aaggaagccca ccctggggc 60
335 gtccacccctt aaatgactgg gcaaacgtt accgactgca tcgggtggatc acggacccgt 120
337 accttcgacg atctgagcga ccgttaccac acccaactcg acccaaggat gaatgcctcc 180
339 cagtccttgg agctcgccctt tatcatcgca gagaggctga ggaagaggag gatcgatcg 240
W--> 341 gggctcaaca gcagccgtcc actgcccgnca ctggctttctt gatgtggccgg agccaaacac 300
343 aaaggggaggtt aggaatagctt gtggtgacctt cggaggagaa gagacagtcg acgccttgc 360
345 tggtgatgc 369
348 <210> SEQ ID NO: 15
349 <211> LENGTH: 362

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Input Set : D:\AprilCornReg.rpt
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350 <212> TYPE: DNA
351 <213> ORGANISM: Zea mays
353 <223> OTHER INFORMATION: Clone ID: LIB3061-002-Q1-K1-D12
355 <400> SEQUENCE: 15
357 gcttaggccta acttgaaggt gggctaaagt ggacaacatg gtggagagcc tttagcggtt 60
359 gtttattecg ccaagactgg gctggattac gtttttgc cccctttag ggtcccgatc 120
361 gcttaggctag ctgcagtc ggtgcgttc tgagggtgc tctgtcttgtt ggcagccgga 180
363 tcgcctgtc ttggtgcatc tggtgatcaa taataactact atgacagacg catgtctgt 240
365 gaagagtatt agtagcagcg ctcataaaaa ctacagttcc gtctctattt tcagctatgt 300
367 aaaaccttcca aactgttcat gcttaaaaact gaaggtttc ttgggtgtgag gtgtgcattgt 360
369 cg 362
372 <210> SEQ ID NO: 16
373 <211> LENGTH: 419
374 <212> TYPE: DNA
375 <213> ORGANISM: Zea mays
377 <223> OTHER INFORMATION: Clone ID: LIB3061-049-Q1-K1-C8
379 <400> SEQUENCE: 16
381 cacccctctcg ctgcactcc gtcgtctga catctcgatc cgtccttcg tttcgaagg 60
383 tcgggagccc cgacgatggc caccaccgcg gtgtcggtcg acgagaagct cgataagctt 120
385 cgcgcggagg tcgcaagct cgacccgatc agcgagaacg agaaggccg gtccatcagc 180
387 ctgcgtcactc ggtacccatc tggggaggcg gagcagatcg agtggagacaa gatccagacc 240
389 ctacccatggat aagtgggtgg gcctctacat acgtccgcg ccgcctccca agatctcgag 300
391 gagacaaga agtgcgttga taagctcgat ttgctcaagc ttaacggagg gtcggagc 360
393 accatgggtt gcatggggcc caagtctgtc attgaagtcc gcaatgggtt cacattct 419
395 <210> SEQ ID NO: 17
397 <211> LENGTH: 431
398 <212> TYPE: DNA
399 <213> ORGANISM: Zea mays
401 <223> OTHER INFORMATION: Clone ID: LIB3061-026-Q1-K1-D7
403 <400> SEQUENCE: 17
405 ctgcgtgtca ctatccgtact tctggaccctt cccctccacg agttcttcc agaaggggggc 60
407 gtcgaggaaa tcgtgcgtga actgtgttct gagacgggag ccaaccagga ggtgcctc 120
409 gcacggatc aaaacgttcc ggaagtaaac cccatgttgc gtttccgtgg ctgcaggctt 180
411 ggttatactgt accctgtatc aacagaaatg caagccgtg ccatctttga agtgcgtata 240
413 cgcgtatccca accaggccgt tcaagttttt ccagatataa tggttccctt tggggaaaca 300
415 ccgcaggaaat tgggacatca ggtggccctt atccgtcaag tcgctaacaa agttttcac 360
417 agtatggggaa aactattggt tacaagatc ggaacgtatc ttgagattcc cagggcagct 420
419 ctatgggtcg a 431
422 <210> SEQ ID NO: 18
423 <211> LENGTH: 319
424 <212> TYPE: DNA
425 <213> ORGANISM: Zea mays
427 <223> OTHER INFORMATION: Clone ID: LIB3061-009-Q1-K1-F12
429 <400> SEQUENCE: 18
431 cagacgcgaa taccctgag gatgcattgg ctgcgcggaa caacggggca gaaggaaatgc 60
433 gggttgtgcc ggacagagca catgttctc gttcggatg agaggattaa ggctgtgagg 120
435 cagatgatca tggccccac agttgtatcg aggacgatcg cgctgtatcg ctttctgcct 180
437 taccagatgt ctgatcttc aggcataatcc cgtgtatgg atggacttcc agtgcgtatc 240
439 cgtacttgcg acccttccact acacqagtcc ctttcaqaag qqaacqtcqa qqaatctcq 300

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Please Note:

Please Note: Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/553,094

DATE: 05/09/2000

TIME: 16:28:07

Input Set : D:\AprilCornReg.rpt
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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:199 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:8
L:199 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:8
L:199 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:8
L:199 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:8
L:212 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:9
L:212 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:9
L:212 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9
L:212 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:9
L:341 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
L:341 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:14
L:341 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:341 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:14
L:488 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:20
L:488 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:20
L:488 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
L:488 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:20
L:861 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:36
L:861 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:36
L:861 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:36
L:861 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:36
L:1090 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:46
L:1090 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:46
L:1090 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:46
L:1090 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:46
L:1143 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:48
L:1143 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:48
L:1143 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:48
L:1143 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:48
L:1290 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:54
L:1290 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:54
L:1290 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:54
L:1290 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:54
L:1405 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:59
L:1405 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:59
L:1405 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:59
L:1405 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:59
L:1422 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:60
L:1422 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:60
L:1422 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:60
L:1422 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:60
L:1455 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:61
L:1455 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:61
L:1455 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:61
L:1455 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:61
L:1578 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:66
L:1578 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:66

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L:1578 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:66
L:1578 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:66
L:1629 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:68
L:1629 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:68
L:1629 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:68
L:1629 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:68
L:1654 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:69
L:1654 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:69
L:1654 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:69
L:1654 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:69
L:1656 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:69
L:1656 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:69
L:1656 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:69
M:340 Repeated in SeqNo=69
L:1683 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:70
L:1683 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:70
L:1683 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:70
L:1683 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:70
L:1776 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:74
L:1776 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:74
L:1776 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:74
L:1919 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:80
M:340 Repeated in SeqNo=80
L:2010 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:84
L:2065 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:86
M:340 Repeated in SeqNo=86
L:2808 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:116
L:2895 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:120
L:2940 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:122
L:3239 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:135
L:3598 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:150
M:340 Repeated in SeqNo=150
L:4291 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:179
L:5294 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:219
L:5403 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:224
L:5428 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:225
L:5649 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:234
L:5700 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:236
L:5725 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:237
L:5796 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:240
L:5813 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:241
L:6274 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:260
L:6343 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:263
L:6370 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:264
L:6539 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:271
L:6568 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:272
L:6615 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:274
L:6672 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:277
L:6749 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:280
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VERIFICATION SUMMARY
PATENT APPLICATION: US/09/553,094

DATE: 05/09/2000
TIME: 16:28:07

Input Set : D:\AprilCornReg.rpt
Output Set: N:\CRF3\05092000\I553094.raw

L:6850 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:284
M:340 Repeated in SeqNo=284
L:6875 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:285
L:7246 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:300
L:7275 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:301